- d) selecting at least one transgenic individual derived from each backcross until the isotransgenic line is produced.
- 2. (AMENDED) The method of Claim 1, wherein the selecting hybrid primary transformants comprises identifying genomic sequences adjacent to the T-DNA inserted and determining the parent genome which has received said T-DNA.
- 3. (AMENDED) The method of Claim 2, wherein determining the plant genome which has received the T-DNA is carried out according to a technique selected from the group consisting of an RFLP technique and a sequencing method.
 - 4. (AMENDED) The method of Claim 1, wherein the individual selected in (d) has
 - a) a chromosome having the T-DNA but otherwise having a geneotype entirely of the line of interest; and
 - b) a genome of interest to entire genome ratio of at least about 75%.
- 5. (AMENDED) The method of Claim 1 further comprising crossing the isotransgenic plant line and a second line of interest.
- 6. (AMENDED) The method of Claim 1, wherein the hybrid plant is selected from the group consisting of a crop plant, vegetable plant, and floral plant.
- 7. (AMENDED) The method of Claim 1, wherein the T-DNA comprises in particular a nucleotide sequence encoding a protein which confers agronomic properties and/or properties of resistance to diseases.
- **8.** (AMENDED) The method of Claim 1, wherein the isotransgenic line is a commercial elite line.
- 9. (AMENDED) The method of Claim 1, wherein the isogenic plant is substantially free of fragments linked to the transgene which may be the subject of a genetic burden.

34934-PCT-USA 072667.0180 PATENT

10. (AMENDED) A method of identifying the parent genome which has received a T-DNA after transformation of a hybrid comprising identifying genomic sequences adjacent to the T-DNA inserted.

- end.
- 11. (AMENDED) A transgenic plant, organ or seed obtained by the method of Claim 1 or 5.
- 12. (AMENDED) An isotransgenic line produced from hybrid transformants by the method of Claims 1 or 7, wherein the isotransgenic line has a pure line of interest genotype over the entire genome and have stably integrated the T-DNA containing the transgene.

Please add the following new claims.

- 14. (NEW) The method of Claim 5, wherein the second line of interest is an isotransgenic plant line.
- 15. (NEW) The method of claim 6, wherein the crop plant or vegatable plant is selected from the group consisting of maize, wheat, rapeseed, sunflower, pea, soybean and barley.
- 16. (NEW) A hybrid according to Claim 12, wherein the hybrid is produced according to the method of claim 5 and wherein the pure line of interest is a commercial crop line.